# Quantification of *Salmonella* spp. using a miniaturisation of MSRV enrichment medium: mini-MSRV

# Quantification of Salmonella spp.

#### Food safety:

retrospective studies of outbreaks: could permit to rapidly identify the highly contaminated part of a control meal

#### Epidemiology

- identify the at risk steps in a primary production chain
- reveal the efficiency of partial decontamination procedure in highly contaminated environment

#### What exists?

- **Direct count :** Immunofluorescent or DNA probes
  - sensibility specificity antibodies (all serovars?)
  - technical steps (filtration...) micro-colonies
  - official method in AOAC and BAM (FDA)

#### • Culture techniques:

- Direct on selective solid medium:
  - selectivity against dominant digestive flora
  - need selective or non selective concentration
  - selective media dedicated (Dulcitol-Bile-Novobiocine)
  - Culturability (stressed Salmonella cells ) overlay procedure,
- Most probable number methods...

# Most probable number methods...

- Theory assumes two conditions:
  - organisms are randomly distributed throughout the solution
  - each sample from the solution which contains at least one organism is able to exhibit growth in the culture medium
- Both hypothesis are false (overall for Salmonella)
  - underline the need to well homogenised the solutions
  - select the most sensitive detection technique.

# Most probable number methods...

- Principe: Based on repetition of serial dilutions of a sample Generate a characteristic number
- Calculation of the MPN:
  - using statistical assumption each characteristic number is related to a MPN with confidence limits
- Increase confidence of the result:
  - increase the number of repetition on the right dilution only if always almost the same level of contamination... so necessity to quantify?

# Most probable number methods...

#### • Limits:

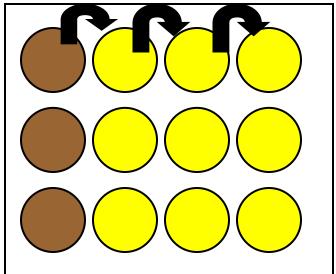
- time and labor consuming
- uncertain results
- if low numbers, do we need quantification?

#### Assuming this we propose

- a rapid and compatible with high number of samples studies (at least 75 samples per week and person)
- based on miniaturised MSRV enrichment
- to discriminated highly from poorly contaminated samples

• Sample diluted 1/10 in EPT: stomaching at least one minute

500μ1 500μ1 500μ1

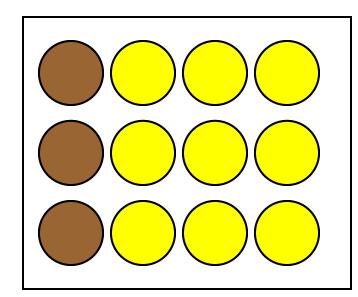


2 mL EPT

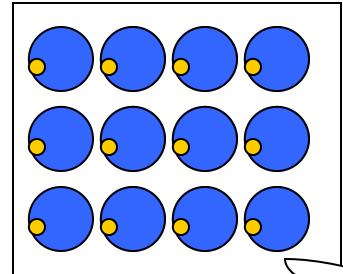
2,5 ml diluted sample

Serial dilution 1/5 with multi-channel pipette

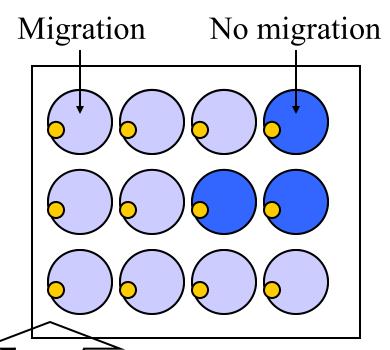
Incubation 16 h 37°C



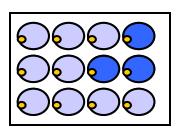
Replication
of the pre-enriched
plate 20µl/well
with multichannel pipette



Incubation 24 - 36h 41,5°C



- Streak each positive well on Rambach agar
- Biochemical characterisation of typical colonies
- Serogroup determination using polyspecific serums
- Determination of the characteristic number of the sample



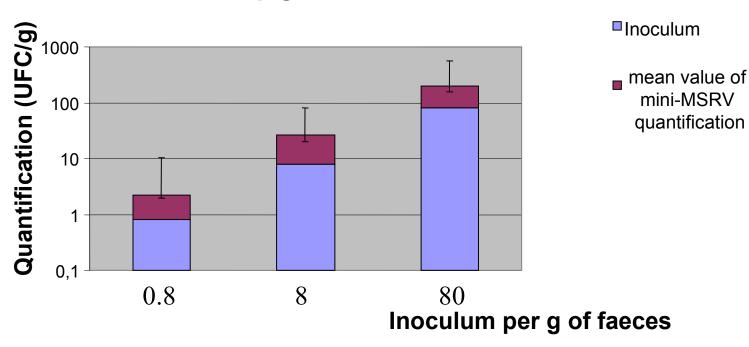
Characteristic number: 3321

http://www.i2workout.com/mcuriale/mpn/index.html MPN deducted :1,33.10<sup>2</sup> (IC :4,910<sup>1</sup>-4.310<sup>2</sup>) *Salmonella* per g of sample

(!Carefull to the initial dilution!)

	mini	Maxi
Characteristic nb	1000	3332
<b>MPN</b> (Salmonella/g)	1,2	$5,7.10^2$
CI	0,17-8,65	$1,5.10^2$ - $2,1.10^3$

# Mean values obtained after mini-MSRV quantification of artificially contaminated pig faeces



N = 7 samples independently inoculated for each inoculum

- Quantification in pig fecal samples
- Quantification in surface samples in lairage
- Quantification in turkey neck skin samples

Quantification in pig fecal samples

	Distribution of samples	Salmonella MPN per g	
	181	0,0	
	25	1,0	
	1	3,5	2
	4	6,4	pigs
	4	3,3.10 <sup>1</sup>	populations
	3	1,8.10 <sup>2</sup> 8,9.10 <sup>3</sup>	
	6	$8,9.10^3$	
TOTAL	224		

2 different risk levels for the production chain

- Quantification in surface samples in lairage
  - 300cm<sup>2</sup> per swab
  - 192 samples, 27 detected by mini-MSRV method
  - 24 positive samples present less than 1 *Salmonella* per cm<sup>2</sup> (MPN=0.82 IC 0.11 5.8)
- Improve the decontamination on homogeneised contaminated surfaces

- Quantification in turkey neck skin samples
  - samples from a positive herd
  - 10 g of neck skin
  - 122 samples, 14 positives, 9 detected by mini-MSRV method

Positive sample	MPN Salmonella /g of skin	IC	
1	0,5	1,4	5,4
2	0,1	0,6	4,2
3	0,1	0,6	4,2
4	1,4	3,9	10,7
5	0,1	0,5	4,2
6	1,3	3,5	9,8
7	0,7	2,1	6,9
8	3,1	8,3	21,9
9	0,1	0,6	4,2

Few numbers of bacteria on the skin of this suspected contaminated batch

#### Conclusion

- First investigations / well defined the need of quantification
- You could adapte the disposition of the plate or the number of plate fonction of the contamination level
- Convenient quantification method for high numbers of samples
- Development: more convenient if we need just to confirm the more diluted positive well